3/26



OIPE

2.

RAW SEQUENCE LISTING

1 <110> APPLICANT: della-Cioppa, Guy

PATENT APPLICATION: US/10/072,438

DATE: 02/28/2002 TIME: 10:29:12

_	\1.1.U>	millional. della eloppa, ea				
2		Erwin, Robert L.				
3		Fitzmaurice, Wayne P.				
4		Hanley, Kathleen M.				
5		Hanley, Kathleen M. Kumagai, Monto H.				
6		Lindbo, John A.				
7		McGee, David R.				
8		Padgett, Hal S.				
9		Poque, Gregory P.				
10	<120>	TITLE OF INVENTION: METHOD OF DETERMINING THE FUNCTION OF				
11		NUCLEOTIDE SEQUENCES AND THE PROTEINS THEY ENCODE BY				
12		TRANSFECTING THE SAME INTO A HOST				
13	<130>	FILE REFERENCE: 08010137US01				
15	<140>	CURRENT APPLICATION NUMBER: 10/072,438				
		CURRENT FILING DATE: 2002-02-05				
18	<150>	PRIOR APPLICATION NUMBER: US/09/232,170				
		PRIOR FILING DATE: 1999-01-15				
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		SEQUENCE: 1				
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35	<212>	TYPE: DNA				
36	<213>	ORGANISM: VIRAL				
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		SEQUENCE: 3				
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RAW SEQUENCE LISTING
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55 <211> LENGTH: 24	
56 <212> TYPE: DNA	
57 <213> ORGANISM: VIRAL	
58 <400> SEQUENCE: 5	
59 tatgtatggt gcagaagaac agat	. 24
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65 <400> SEQUENCE: 6	
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68 <210> SEQ ID NO: 7	
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70 <212> TYPE: DNA	
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75 <210> SEQ ID NO: 8	
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78 <213> ORGANISM: VIRAL	
79 <400> SEQUENCE: 8	. 20
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89 <210> SEQ ID NO: 10	
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91 <212> TYPE: DNA	
92 <213> ORGANISM: VIRAL	
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97 <211> LENGTH: 44 98 <212> TYPE: DNA	
99 <213> ORGANISM: VIRAL	
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105 <212> TYPE: DNA	
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108 aagatetete gagetaaaeg ggaegetgee aaagacegge egg	43
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124	<210>	SEQ ID NO: 15			
125	<211>	LENGTH: 32			
126	<212>	TYPE: DNA			
127	<213>	ORGANISM: VIRAL			
128	<400>	SEQUENCE: 15			2.2
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131	<210>	SEQ ID NO: 16			
		LENGTH: 31			
		TYPE: DNA			
134	<213>	ORGANISM: VIRAL			
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	Z2105	SEQ ID NO: 18	,		
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		SEQUENCE: 18			
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		LENGTH: 36			
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				•	
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164	40 1 O		aagaccygcc	200209	
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168	<212>	TYPE: DNA	
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192		tccctaggct aaatagcata acttccacat caaagc	36
194	<210>	SEQ ID NO: 25	
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201	<210>	SEQ ID NO: 26	
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217	<212>	TYPE: DNA	
218	<213>	ORGANISM: VIRAL	
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		SEQ ID NO: 29	
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/072,438 TIME: 10:29:12

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	11> LENGTH: 36					
	12> TYPE: DNA 13> ORGANISM: VIRAL					
	00> SEQUENCE: 30	•				
233 < 4	cqgqgtacct gggcccctac	ccaaaattta	aaaaaa			36
	10> SEQ ID NO: 31	0099990004	555~55			
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239 <2	13> ORGANISM: VIRAL	•				
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241	ctcgaggttc ataagaccgc					60
242	ttgtcagata aaaggttgtt					120
243	gcctaaagtt agtgacttcc					180
244	gacgagatta aagactgtct					240
245	gtccctgtgt gatattgatt					300
246	tgttttgggt gttgttttca					360
247	cagtgagcgt gattgacaaa					420
248	gagctgctgt aaaggacaga					480 540
249	ttgcggatgc caagcgaaaa					600
250	agctggatgg caacctctag					660
251 252	ggttgttaaa ggtttgaggg gaaggtgtgg ttgacgattt					720
252	cgaaagaaaa aagaaaaaga					780
253 254	accggagaga tacgccggtc					840
255	tcacgagete gaatcagtac					900
256	ttaaccagtg cgtgtctgcg	ttgtcgcaat	cgtatcaaac	t.caggcggca	agagatactg	960
257	ttagacagca gttctctaac					1020
258	aaacaggata ccgggtgtat					1080
259	tgaagteett tgataetaga					1140
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261	caaattcagc ttttgctgaa	cgagctctcc	aacggacatg	gtctgatcga	acagggcaga	1260
262	gttcgaggtt ttattacctt	gggctactgc	gccagctaca	taggcgtggt	gcacacgata	1320
263	gtgcatagtg tttttctctc					1380
264	agggtggcgt aaaccaaatt					1440
265	ttcctggatc acctgttaac		_	agtgggaata	actaaaagtg	1500
266	agagttegaa teeteetaae	cccgggtagg	ggccca			1536
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	12> TYPE: DNA					
	13> ORGANISM: VIRAL					
272 <4 273	00> SEQUENCE: 32 gatggcgcct taatacgacg	gagtatagtt	ttattttat	tacaacaaca	acaac	55
	10> SEQ ID NO: 33	cactataget	ccacccccyc	cycaacaca	acaac	,,,
	10> SEQ 1D NO. 33 11> LENGTH: 30					
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/072,438

DATE: 02/28/2002 TIME: 10:29:13

Input Set : N:\Crf3\RULE60\10072438.raw.txt
Output Set: N:\CRF3\02282002\J072438.raw

:369 M:351 W: Sequence data Name/Key Feature Out-of-Range, SEQ ID#:43, CDS LOCATION: (1)... 1992) 3:373 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 3:376 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 3:379 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 :382 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 :385 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 :388 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 .:391 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 :394 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 J:397 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 J:400 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 :403 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 3:406 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 :409 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 :412 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 3:415 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43 J:418 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:43